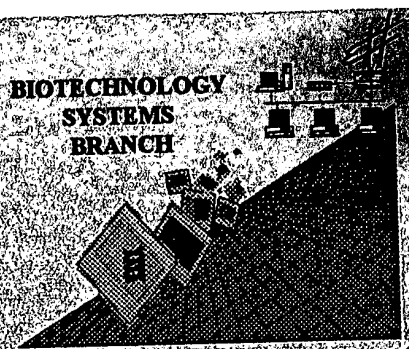


Fitzgerald

BIOTECHNOLOGY
SYSTEMS
BRANCH

RAW SEQUENCE LISTING ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following CRF diskette:

Application Serial Number:

09/029,042A

Art Unit / Team No.

1646

Date Processed by STIC:

10/28/99

THE ATTACHED PRINTOUT EXPLAINS THE ERRORS DETECTED.

PLEASE BE SURE TO FORWARD THIS INFORMATION TO THE APPLICANTS BY EITHER:

1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANTS ALONG WITH A NOTICE TO COMPLY or,

2) CALLING APPLICANTS AND FAXING THEM A COPY OF THE PRINTOUT WITH A NOTICE TO COMPLY

THIS WILL INSURE THAT THE NEXT SUBMISSION RECEIVED FROM THEM WILL BE ERROR FREE.

IF YOU HAVE ANY FURTHER QUESTIONS, PLEASE CALL:

MARK SPENCER 703-308-4212

Raw Sequence Listing Error Summary

ERROR DETECTED SUGGESTED CORRECTION

SERIAL NUMBER: 09/029,042A

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 Wrapped Nucleics The number/text at the end of each line "wrapped" down to the next line.
This may occur if your file was retrieved in a word processor after creating it.
Please adjust your right margin to .3, as this will prevent "wrapping".
- 2 Wrapped Aminos The amino acid number/text at the end of each line "wrapped" down to the next line.
This may occur if your file was retrieved in a word processor after creating it.
Please adjust your right margin to .3, as this will prevent "wrapping".
- 3 Incorrect Line Length The rules require that a line not exceed 72 characters in length. This includes spaces.
- 4 Misaligned Amino Acid Numbering The numbering under each 5th amino acid is misaligned. This may be caused by the use of tabs between the numbering. It is recommended to delete any tabs and use spacing between the numbers.
- 5 Non-ASCII This file was not saved in ASCII (DOS) text, as required by the Sequence Rules.
Please ensure your subsequent submission is saved in ASCII text so that it can be processed.
- 6 Variable Length Sequence(s) contain n's or Xaa's which represented more than one residue.
As per the rules, each n or Xaa can only represent a single residue.
Please present the maximum number of each residue having variable length and indicate in the (ix) feature section that some may be missing.
- 7 PatentIn ver. 2.0 "bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequence(s) . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence.
- 8 Skipped Sequences (OLD RULES) Sequence(s) missing. If intentional, please use the following format for each skipped sequence:
(2) INFORMATION FOR SEQ ID NO:X:
(i) SEQUENCE CHARACTERISTICS:(Do not insert any headings under "SEQUENCE CHARACTERISTICS")
(xl) SEQUENCE DESCRIPTION:SEQ ID NO:X:
This sequence is intentionally skipped

Please also adjust the "(iii) NUMBER OF SEQUENCES:" response to include the skipped sequence(s).
- 9 Skipped Sequences (NEW RULES) Sequence(s) missing. If intentional, please use the following format for each skipped sequence.
<210> sequence id number
<400> sequence id number
000
- 10 Use of n's or Xaa's (NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.
Use of <220> to <223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
- 11 Use of <213>Organism (NEW RULES) Sequence(s) are missing this mandatory field or its response.
- 12 Use of <220>Feature (NEW RULES) Sequence(s) are missing the <220>Feature and associated headings.
Use of <220> to <223> is MANDATORY if <213>ORGANISM is "Artificial" or "Unknown"
Please explain source of genetic material in <220> to <223> section.
(See "Federal Register," 6/01/98, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of new Rules)
- 13 PatentIn ver. 2.0 "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing).
Instead, please use "File Manager" or any other means to copy file to floppy disk.

PAGE: 1

RAW SEQUENCE LISTING

PATENT APPLICATION US/09/029,042A

 DATE: 10/28/1999
 TIME: 14:00:56

Input Set: I029042A.RAW

This Raw Listing contains the General Information Section and those Sequences containing ERRORS.

Does Not Comply
Corrected Diskette Needed

- 1 <110> Kim, Sun-Young; Kim, Kee-Won; Kim, Tae-Han; Hwang, Jeong-Ho; Kim, Seon-Hee; L
 2 <120> Heterologous Protein Production System using Avian Cells
 3 <130>
 4 <140> US/09/029,042A
 5 <141> 1998-05-15
 6 <150> PCT/KR96/00145
 7 <151> 1996-08-23
 8 <160> 11

W--> OK

Please consult
new sequence Rules
and sample sequence
Listing (attached in back)
for valid format.

all text
must be
visible -
see item 3
on Error
summary sheet

ERRORED SEQUENCES FOLLOW

insert sequence number NEXT TO <400>

9 <210> 1
 10 <211> 1584
 11 <212> DNA
 12 <213> erythropoietin
 13 <400> ATGGGGGTGC ACGAATGTCC TGCCTGGCTG TGGCTTCTCC TGTCCTGTCT
 14 GTCGCTCCCT CTGGGCCTCC CAGTCTGGG CGCCCCACCA CGCCTCATCT
 15 GTGACAGCCG AGTCTCTGGAG AGGTACCTCT TGGAGGCCAA GGAGGCCGAG
 16 AATATCACGG TGAGACCCCT TCCCCAGCAC ATTCCACAGA ACTCACGCTC
 17 AGGGCTTCAG GG-AACTCCT CCCAG-ATCC AGGAACCTGG CACTTGGTTT
 18 GGGGTGGAGT TGGGAAGCTA GACACTGCCC CCCTACATAA GAATAAGTCT
 19 GGTGGCCCCA AACCATACCT GGAAACTAGG CAAGGAGCAA AGCCAGCAGA
 20 TCCTACGGCC TGTGGGCCAG GGCCAG-AGC CTTACAGGAC CTTGACTCC
 21 CCGGGCTGTG TGCATTTTCTG ACGGGCTGTG CTGAACACTG CAGCTTGAAT
 22 GAGAATATCA CTGTCCCGA CACCAAAGTT AATTTCTATG CCTGGAAGAG
 23 GATGGAGGTG AGTTCCTTTT TTTTTTTTTT TCCTTTCTTT TGGAGAATCT
 24 CATTTGCGAG CCTGATTTTG GATGAAAGGG AGAATGATCG GGGGAAAGGT
 25 AAAATGGAGC AGCAGAGATG AGGCTGCCTG GGCGCAGAGG CTCACGTCTA
 26 TAATCCCAGG CTGAGATGGC CGAGATGGGA GAATTGCTTG AGCCCTGGAG

DO NOT USE <400> line

So
100
↓

Per 1.822
of sequence
Rules, insert
the cumulative
base total at
end of each
line.

DO NOT use
upper-case
letters for
nucleic acids;
under new
sequence Rules,
use lower-case
letters for bases.

PAGE: 2

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/029,042A

DATE: 10/28/1999

TIME: 14:00:56

Input Set: I029042A.RAW

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*Insert
Cumulative
base totals*

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79 <213> erythropoietin 2

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same error

PAGE: 3

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/029,042ADATE: 10/28/1999
TIME: 14:00:56

Input Set: I029042A.RAW

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E--> 137     CCGCAAATC TTCCGAGTCT ACTCCAATT CCTCCGGGGA AAGCTGAAGC
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PAGE: 4

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/029,042ADATE: 10/28/1999
TIME: 14:00:56

Input Set: I029042A.RAW

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PAGE: 5

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/029,042A

DATE: 10/28/1999

TIME: 14:00:56

Input Set: I029042A.RAW

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211 <213> erythropoietin -> 4
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W--> 218 AGGGCTTCAG GGAACTCCT CCCAGATCC AGGAACCTGG CACTTGGTTT
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PAGE: 6

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/029,042A

DATE: 10/28/1999
TIME: 14:00:56

Input Set: I029042A.RAW

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PAGE: 7

RAW SEQUENCE LISTING
 PATENT APPLICATION US/09/029,042A

DATE: 10/28/1999

TIME: 14:00:56

Input Set: I029042A.RAW

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PAGE: 8

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/029,042ADATE: 10/28/1999
TIME: 14:00:56

Input Set: I029042A.RAW

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W--> 372 GTTCAGACCA ACCTAGGCAG CCTAGTGAGA TCCCCATCT CTACAAACAT
E--> 373
W--> 374 TTAAAAAAT TAGTCAGGTG AAGTGGTGCA TGGTGGTAGT CCCAGATATT
E--> 375
W--> 376 TGGAAGGCTG AGGCGGGAGG ATCGCTTGAG CCCAGGAATT TGAGGCTGCA
E--> 377
W--> 378 GTGAGCTGTG ATCACACCAC TGCCTCCAG CCTCAGTGAC AGAGTGAGGC
E--> 379
W--> 380 CCTGTCTCAA AAAAGAAAAG AAAAAAGAAA AATTATGAGG GCTGTATGGA
E--> 381
W--> 382 ATACATTCAT TATTCATTCA CTCCTCACT CACTCATTCA TTCATTCATT
E--> 383
W--> 384

PAGE: 9

RAW SEQUENCE LISTING PATENT APPLICATION US/09/029,042A

DATE: 10/28/1999
TIME: 14:00:56

Input Set: I029042A.RAW

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E--> 385      CATTCAACAA GTCTTATTGC ATACCTTCTG TTTGCTCAGC TTGGTGCTTG
W--> 386      GGGCTGCTGA GGGGCAGGAG GGAGAGGGTG ACATGGGITC ACTGACTCCC
E--> 387
W--> 388      AGAGTCCACT CCCTGTAGGT CGGGCAGCAG GCCGTAGAAG TCTGGCAGGG
E--> 389
W--> 390      CCTGGCCCTG CTGTCGGAAG CTGTCCTGCG GGGCCAGGCC CTGTTGGTCA
E--> 391
W--> 392      ACTCTTCCCA GCCGTGGGAG CCCCTGCAGC TGCATGTGGA TAAAGCCGTC
E--> 393
W--> 394      AGTGGCCTTC GCAGCCTCAC CACTCTGCTT CGGGCTCTGG GAGCCCAGGT
E--> 395
W--> 396      GAGTAGGAGC GGACACTTCT GCTTGCCCTT TCTGTAAGAA GGGGAGAAGG
E--> 397
W--> 398      GTCTTGCTAA GGAGTACAGG ATCTGTCCGT ATTCCTTCCC TTTCTGTGGC
E--> 399
W--> 400      ACTGCAGCGA CCACCTGTTT TCTCCTTGGC AGAAGGAAGC CATCTCCCCT
E--> 401
W--> 402      CCAGATGCGG CCTCAGCTGC TCCACTCCGA ACAATCACTG CTGACACTTT
E--> 403
W--> 404      CCGCAAATC TTCCGAGTCT ACTCCAATTT CCTCCGGGGA GAGCTGAAGC
E--> 405
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E--> 407
W--> 408

```

```

E--> 409      <210> 7
E--> 410      <211> 193
E--> 411      <212> PRT
E--> 412      <213> erythropoietin gene
E--> 413      <400> Met Gly Val His Glu Cys Pro Ala Trp Leu Trp Leu Leu Ser Leu
E--> 414      Leu Ser Leu Pro Leu Gly Leu Pro Val Leu Gly Ala Pro Arg Leu
E--> 415      Ile Cys Asp Ser Arg Val Leu Glu Arg Tyr Leu Leu Glu Ala Lys Glu
E--> 416      Ala Glu
E--> 417      Asn Ile Thr Thr Gly Cys Ala Glu His Cys Ser Leu Asn Glu Asn Ile
E--> 418      Thr Val Pro Asp Thr Lys Val Asn Phe Tyr Ala Trp Lys Arg Met Glu
E--> 419      Val Gly Gln Gln Ala Val Glu Val Trp Gln Gly Leu Ala Leu Leu Ser
E--> 420      Glu Ala
E--> 421      Val Leu Arg Gly Gln Ala Leu Leu Val Asn Ser Ser Gln Pro Trp Glu
E--> 422      Pro Leu Gln Leu His Val Asp Lys Ala Val Ser Gly Leu Arg Ser Leu
E--> 423      Thr Thr Leu Leu Arg Ala Leu Gly Ala Gln Lys Glu Ala Ile Ser Pro
E--> 424
E--> 425
E--> 426
E--> 427
E--> 428
E--> 429
E--> 430
E--> 431
E--> 432
E--> 433

```

do not show amino acids next to
24007

per 1.822

of
sequence
number

the amino
acid
order

every 5
amino acids.

Do NOT use
TAB codes

between
amino acid nos.

Use space
characters.

PAGE: 10

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/029,042A

DATE: 10/28/1999
TIME: 14:00:56

Input Set: I029042A.RAW

434 Pro Asp
E--> 435 Ala Ala Ser Ala Ala Pro Leu Arg Thr Ile Thr Ala Asp Thr Phe Arg
436
E--> 437 Lys Leu Phe Arg Val Val Ser Asn Phe Leu Arg Gly Lys Leu Lys Leu
438
E--> 439 Tyr Thr Gly Glu Ala Cys Arg Thr Gly Asp Arg
440
E--> 441

442 <210> 8
E--OK 443 <211> 193
444 <212> PRT
445 <213> erythropoietin gene
E--> 446 <400> Met Gly Val His Glu Cys Pro Ala Trp Leu Trp Leu Leu Leu Ser Leu
447 Leu Ser Leu Pro Leu Gly Leu Pro Val Leu Gly Ala Pro Pro Arg Leu
E--> 448 Ile Cys Asp Ser Arg Val Leu Glu Arg Tyr Leu Leu Glu Ala Lys Glu
449
E--> 450 Ala Glu
451
E--> 452 Asn Ile Thr Thr Gly Cys Ala Glu His Cys Ser Leu Asn Glu Asn Ile
453
E--> 454 Thr Val Pro Asp Thr Lys Val Asn Phe Tyr Ala Trp Lys Arg Met Glu
455
E--> 456 Val Gly Gln Gln Ala Val Glu Val Trp Gln Gly Leu Ala Leu Leu Ser
457
E--> 458 Glu Ala
459
E--> 460 Val Leu Arg Gly Gln Ala Leu Leu Val Asn Ser Ser Gln Pro Trp Glu
461
E--> 462 Pro Leu Gln Leu His Val Asp Lys Ala Val Ser Gly Leu Arg Ser Leu
463
E--> 464 Thr Thr Leu Leu Arg Ala Leu Gly Ala Gln Lys Glu Ala Ile Ser Pro
465
E--> 466 Pro Asp
467
E--> 468 Ala Ala Ser Ala Ala Pro Leu Arg Thr Ile Thr Ala Asp Thr Phe Arg
469
E--> 470 Lys Leu Phe Arg Val Val Ser Asn Phe Leu Arg Gly Lys Leu Lys Leu
471
E--> 472 Tyr Thr Gly Glu Ala Cys Arg Thr Gly Asp Arg
473
E--> 474

475 <210> 9
E--OK 476 <211> 193
477 <212> PRT
478 <213> erythropoietin gene
E--> 479 <400> Met Gly Val His Glu Cys Pro Ala Trp Leu Trp Leu Leu Leu Ser Leu
480 Leu Ser Leu Pro Leu Gly Leu Pro Val Leu Gly Ala Pro Pro Arg Leu
E--> 481

same error

number
the
amino
acids
under
every 5
amino
acids
Do NOT
use TAB
codes.

number
amino
acids

PAGE: 11

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/029,042A

DATE: 10/28/1999
TIME: 14:00:56

Input Set: I029042A.RAW

482 Ile Cys Asp Ser Arg Val Leu Glu Arg Tyr Leu Leu Glu Ala Lys Glu
E--> 483 Ala Glu
484
E--> 485 Asn Ile Thr Lys Gly Cys Ala Glu His Cys Ser Leu Asn Glu Asn Ile
486
E--> 487 Thr Val Pro Asp Thr Lys Val Asn Phe Tyr Ala Trp Lys Arg Met Glu
488
E--> 489 Val Gly Gln Gln Ala Val Glu Val Trp Gln Gly Leu Ala Leu Leu Ser
490
E--> 491 Glu Ala
492
E--> 493 Val Leu Arg Gly Gln Ala Leu Leu Val Asn Ser Ser Gln Pro Trp Glu
494
E--> 495 Pro Leu Gln Leu His Val Asp Lys Ala Val Ser Gly Leu Arg Ser Leu
496
E--> 497 Thr Thr Leu Leu Arg Ala Leu Gly Ala Gln Lys Glu Ala Ile Ser Pro
498
E--> 499 Pro Asp
500
E--> 501 Ala Ala Ser Ala Ala Pro Leu Arg Thr Ile Thr Ala Asp Thr Phe Arg
502
E--> 503 Lys Leu Phe Arg Val Tyr Ser Asn Phe Leu Arg Gly Lys Leu Lys Leu
504
E--> 505 Tyr Thr Gly Glu Ala Cys Arg Thr Gly Asp Arg
506

507 <210> 10
E--OK 508 <211> 193
509 <212> PRT
510 <213> erythropoietin gene -> 10
E--> 511 <400> Met Gly Val His Glu Cys Pro Ala Trp Leu Trp Leu Leu Leu Ser Leu
512 Leu Ser Leu Pro Leu Gly Leu Pro Val Leu Gly Ala Pro Pro Arg Leu
E--> 513 Ile Cys Asp Arg Arg Val Leu Glu Arg Tyr Leu Leu Glu Ala Lys Glu
514
E--> 515 Ala Glu
516
E--> 517 Asn Ile Thr Thr Gly Cys Ala Glu His Cys Ser Leu Asn Glu Asn Ile
518
E--> 519 Thr Val Pro Asp Thr Lys Val Asn Phe Tyr Ala Trp Lys Arg Met Glu
520
E--> 521 Val Gly Gln Gln Ala Val Glu Val Trp Gln Gly Leu Ala Leu Leu Ser
522
E--> 523 Glu Ser
524
E--> 525 Val Leu Arg Gly Gln Ala Leu Leu Val Asn Ser Ser Gln Pro Trp Glu
526
E--> 527 Pro Leu Gln Leu His Val Asp Lys Ala Val Ser Gly Leu Arg Ser Leu
528
E--> 529 Thr Thr Leu Leu Arg Ala Leu Gly Ala Gln Lys Glu Ala Ile Ser Pro
530

PAGE: 12

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/029,042A

DATE: 10/28/1999
TIME: 14:00:56

Input Set: I029042A.RAW

E--> 531
532 Pro Asp
E--> 533
534 Ala Ala Ser Ala Ala Pro Leu Arg Thr Ile Thr Ala Asp Thr Phe Arg
E--> 535
536 Lys Leu Phe Arg Val Tyr Ser Asn Phe Leu Arg Gly Lys Leu Lys Leu
E--> 537
538 Tyr Thr Gly Glu Ala Cys Arg Thr Gly Asp Arg
E--> 539

540 <210> 11
E--> OK 541 <211> 193
542 <212> PRT
543 <213> erythropoietin gene > //
E--> 544 <400> Met Gly Val His Glu Cys Pro Ala Trp Leu Trp Leu Leu Leu Ser Leu
545 Leu Ser Leu Pro Leu Gly Leu Pro Val Leu Gly Ala Pro Pro Arg Leu
E--> 546
547 Ile Cys Asp Ser Arg Val Leu Glu Arg Tyr Leu Leu Glu Ala Lys Glu
E--> 548
549 Ala Glu
E--> 550
551 Asn Ile Thr Thr Gly Cys Ala Glu His Cys Ser Leu Asn Glu Asn Ile
E--> 552
553 Thr Val Pro Asp Thr Lys Val Asn Phe Tyr Ala Trp Lys Arg Met Glu
E--> 554
555 Val Gly Gln Gln Ala Val Glu Val Trp Gln Gly Leu Ala Leu Leu Ser
E--> 556
557 Glu Ala
E--> 558
559 Val Leu Arg Gly Gln Ala Leu Leu Val Asn Ser Ser Gln Pro Trp Glu
E--> 560
561 Pro Leu Gln Leu His Val Asp Lys Ala Val Ser Gly Leu Arg Ser Leu
E--> 562
563 Thr Thr Leu Leu Arg Ala Leu Gly Ala Gln Lys Glu Ala Ile Ser Pro
E--> 564
565 Pro Asp
E--> 566
567 Ala Ala Ser Ala Ala Pro Leu Arg Thr Ile Thr Ala Asp Thr Phe Arg
E--> 568
569 Lys Leu Phe Arg Val Tyr Ser Asn Phe Leu Arg Gly Glu Leu Lys Leu
E--> 570
571 Tyr Thr Gly Glu Ala Cys Arg Thr Gly Asp Gly
E--> 572

1
Delete at end of file

1 FYI

Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the <220> to
<223> fields of each sequence which presents at least one n or Xaa.

Appendix A To Subpart G to Part 1—Sample Sequence Listing

<110> Smith, John

Smith, Jane

<120> Example of a Sequence Listing

<130> 01-00001

<140> US 08/999,999

<141> 1998-02-28

<150> EP 91000000

<151> 1997-12-31

Please consult.

<160> 2

<170> PatentIn ver. 2.0

<210> 1

<211> 403

<212> DNA

<213> Paramecium aurelia

<220>

<221> CDS

<222> 341..394

<300>

<301> Doe, Richard

<302> Isolation and Characterization of a Gene Encoding a

Protease from Paramecium sp.

<303> Journal of Fictional Genes

<304> 1

<305> 4

<306> 1 - 7

<307> 1988-06-20

<400> 1

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ctgactgact ctgagatagt cgagcccgta cgagaccgt cgagggtgac agagagtggg 180

cgcggtgcgcg cagagcgccg cgccgggtgcg cgcgcgagtg cgcggtgggc cgcgcgaggg 240

ctttcgcggc agcggcggcg ctttcggcg cgcgcccgtc cgcccctaga cctgagaggt 300

cttctcttcc ctctcttca ctagagaggt ctatatatac atg gtt tca atg ttc 355

Met Val Ser Met Phe

agc ttg tct ttc aaa tgg cct gga ttt tgt ttg ttt gtt tgtttgctc 403

Ser Leu Ser Phe Lys Trp Pro Gly Phe Cys Leu Phe Val

10

15

<210> 2

<211> 18

<212> PRT

<213> Paramecium aurelia

<400> 2

Met Val Ser Met Phe Ser Leu Ser Phe Lys Trp Pro Gly Phe Cys Leu

1

5

10

15

Phe Val

ed: May 22, 1998.

A. Lehman,

ant Secretary of Commerce and
missioner of Patents and Trademarks.

oc. 98-14194 Filed 5-29-98; 8:45 am]

1 CODE 3510-16-C

identifiers and their accompanying information as shown in the following table. The numeric identifier shall be used only in the "Sequence Listing." The order and presentation of the items of information in the "Sequence Listing" shall conform to the arrangement given below. Each item of information shall begin on a new line and shall begin with the numeric identifier enclosed in angle brackets as shown. The submission of those items of information designated with an "M" is mandatory. The submission of those items of information designated with an "O" is optional. Numeric identifiers <110> through <170> shall only be set forth at the beginning of the "Sequence Listing." The following table illustrates the numeric identifiers.

Numeric Identifier	Definition	Comments and Format	Mandatory (M) or Optional (O)
<110>	Applicant	Preferably max. of 10 names; one name per line; preferable format: Surname, Other Names and/or Initials	M
<120>	Title of Invention		M
<130>	File Reference	Personal file reference	M when filed prior to assignment of appl. number
<140>	Current Application Number	Specify as: US 07/999,999 or PCT/US96/99999	M, if available
<141>	Current Filing Date	Specify as: yyyy-mm-dd	M, if available
<150>	Prior Application Number	Specify as: US 07/999,999 or PCT/US96/99999	M, if applicable include priority documents under 35 USC 119 and 120
<151>	Prior Application Filing Date	Specify as: yyyy-mm-dd	M, if applicable
<160>	Number of SEQ ID NOs	Count includes total number of SEQ ID NOs	M
<170>	Software	Name of software used to create the Sequence Listing	O
<210>	SEQ ID NO: #:	Response shall be an integer representing the SEQ ID NO shown	M
<211>	Length	Respond with an integer expressing the number of bases or amino acid residues	M

<212>	Type	Whether presented sequence molecule is DNA, RNA, or PRT (protein). If a nucleotide sequence contains both DNA and RNA fragments, the type shall be "DNA." In addition, the combined DNA/RNA molecule shall be further described in the <220> to <223> feature section.	M
<213>	Organism	Scientific name, i.e. Genus/species, Unknown or Artificial Sequence. In addition, the "Unknown" or "Artificial Sequence" organisms shall be further described in the <220> to <223> feature section.	M
<220>	Feature	Leave blank after <220>. <221-223> provide for a description of points of biological significance in the sequence.	M, under the following conditions: if "n," "Xaa," or a modified or unusual L-amino acid or modified base was used in a sequence; if ORGANISM is "Artificial Sequence" or "Unknown"; if molecule is combined DNA/RNA.
<221>	Name/Key	Provide appropriate identifier for feature, preferably from WIPO Standard ST.25 (1998), Appendix 2, Tables 5 and 6	M, under the following conditions: if "n," "Xaa," or a modified or unusual L-amino acid or modified base was used in a sequence
<222>	Location	Specify location within sequence; where appropriate state number of first and last bases/amino acids	M, under the following conditions: if "n," "Xaa," or a modified or unusual L-amino acid or modified

		in feature	base was used in a sequence
<223>	Other Information	Other relevant information; four lines maximum	M, under the following conditions: if "n," "Xaa," or a modified or unusual L-amino acid or modified base was used in a sequence; if ORGANISM is "Artificial Sequence" or "Unknown"; if molecule is combined DNA/RNA.
<300>	Publication Information	Leave blank after <300>	0
<301>	Authors	Preferably max of ten named authors of publication; specify one name per line; preferable format: Surname, Other Names and/or Initials	0
<302>	Title		0
<303>	Journal		0
<304>	Volume		0
<305>	Issue		0
<306>	Pages		0
<307>	Date	Journal date on which data published; specify as yyyy-mm-dd, MMM-yyyy or Season-yyyy	0
<308>	Database Accession Number	Accession number assigned by database including database name	0
<309>	Database Entry Date	Date of entry in database; specify as yyyy-mm-dd or MMM-yyyy	0
<310>	Patent Document Number	Document number; for patent-type citations only. Specify as, for example, US 07/999,999	0

<311>	Patent Filing Date	Document filing date, for patent-type citations only; specify as yyyy-mm-dd	O
<312>	Publication Date	Document publication date, for patent-type citations only; specify as yyyy-mm-dd	O
<313>	Relevant Residues	FROM (position) TO (position)	O
<400>	Sequence	SEQ ID NO should follow the numeric identifier and should appear on the line preceding the actual sequence	M

5. Section 1.824 is revised to read as follows:

1.824 Form and format for nucleotide and/or amino acid sequence submissions in computer readable form.

(a) The computer readable form required by 1.821(e) shall meet the following specifications:

(1) The computer readable form shall contain a single "Sequence Listing" as either a diskette, series of diskettes, or other permissible media outlined in paragraph (c) of this section.

(2) The "Sequence Listing" in paragraph (a) (1) of this section shall be submitted in American Standard Code for Information Interchange (ASCII) text. No other formats shall be allowed.

(3) The computer readable form may be created by any means, such as word processors, nucleotide/amino acid sequence editors or other custom computer programs; however, it shall conform to all specifications detailed in this section.

(4) File compression is acceptable when using diskette media, so long as the compressed file is in a self-extracting format that will decompress on one of the systems described in paragraph (b) of this section.

(5) Page numbering shall not appear within the computer readable form version of the "Sequence Listing" file.

(6) All computer readable forms shall have a label permanently affixed thereto on which has been hand-printed or typed: the name of the applicant, the title of the invention, the date on which the data were recorded on the computer readable form, the operating system used, a reference number, and an application serial number and filing date, if known.

(b) Computer readable form submissions must meet these format requirements:

(1) Computer: IBM PC/XT/AT, or compatibles, or Apple Macintosh;

(2) Operating System: MS-DOS, Unix or Macintosh;